

5. A leptin receptor (OB-R) polypeptide which is selected from the group consisting of OB-Ra (SEQ ID NO:2), OB-Rb (SEQ ID NO:4), OB-Rc (SEQ ID NO:6), OB-Rd (SEQ ID NO:8), and OB-Re (SEQ ID NO:10), or allelic variants thereof.
6. A leptin receptor (OB-R) polypeptide which is selected from the group consisting of:
- a) N-terminal corresponding to OB-Ra through Lys<sup>889</sup> and C-terminal to a C-terminal selected from the group consisting of OB-Rb after Lys<sup>889</sup> (SEQ ID NO:57), OB-Rc after Lys<sup>889</sup> (SEQ ID NO:58), and OB-Rd after Lys<sup>889</sup> (SEQ ID NO:59);
  - b) N-terminal corresponding to OB-Rb or OB-Rc through Lys<sup>889</sup>, and C-terminal corresponding to OB-Ra after Lys<sup>889</sup> (SEQ ID NO:60,61) or OB-Rd after Lys<sup>889</sup> (SEQ ID NO:62,63);
  - c) N-terminal corresponding to OB-Rd through Lys<sup>889</sup>, and C-terminal corresponding to OB-Ra after Lys<sup>889</sup> (SEQ ID NO:64), OB-Rb after Lys<sup>889</sup> (SEQ ID NO:65), or OB-Rc after Lys<sup>889</sup> (SEQ ID NO:66);
  - d) N-terminal corresponding to SEQ ID NO:55 from Pro<sup>664</sup> to Lys<sup>889</sup>, and C-terminal corresponding to OB-Ra after Lys<sup>889</sup> (SEQ ID NO:67), OB-Rb after Lys<sup>889</sup> (SEQ ID NO:68), OB-Rc after Lys<sup>889</sup> (SEQ ID NO:69), or OB-Rd after Lys<sup>889</sup> (SEQ ID NO:70);
  - e) N-terminal corresponding to SEQ ID NO:55 from Met<sup>733</sup> to Lys<sup>889</sup>, and C-terminal corresponding to OB-Ra after Lys<sup>889</sup> (SEQ ID NO:71), OB-Rb after Lys<sup>889</sup> (SEQ ID NO:72), OB-Rc after Lys<sup>889</sup> (SEQ ID NO:73), or OB-Rd after Lys<sup>889</sup> (SEQ ID NO:74);
  - f) N-terminal selected from the group consisting of OB-Ra, OB-Rb, OB-Rd, and OB-R from Pro<sup>664</sup> to His<sup>796</sup>, and OB-Re from His<sup>796</sup> (SEQ ID NO:75, 76, 77, and 78);
  - g) N-terminal corresponding to SEQ ID NO:55 from Met<sup>733</sup> to His<sup>796</sup>, and OB-Re from His<sup>796</sup> (SEQ ID NO:79); and
  - h) allelic variants of any of subparts a) through g).

7. A leptin receptor (OB-R) polypeptide wherein

- C1
- a) the N-terminal sequence is selected from the group consisting of
    - i. amino acid residues 1-889 (SEQ ID NO:80);
    - ii. amino acid residues 23-889 (SEQ ID NO:81);
    - iii. amino acid residues 28-889 (SEQ ID NO:82);
    - iv. amino acid residues 133-889 (SEQ ID NO:83);
    - v. amino acid residues 733-889 (SEQ ID NO:84);
    - vi. amino acid residues 1-796 (SEQ ID NO:85);
    - vii. amino acid residues 23-796 (SEQ ID NO:86);
    - viii. amino acid residues 28-796 (SEQ ID NO:87);
    - ix. amino acid residues 133-796 (SEQ ID NO:88);
    - x. amino acid residues 733-796 (SEQ ID NO:89); and
    - xi) allelic variants of any of subparts i) through x); and
  - b) the C-terminal sequence is selected from the group consisting of
    - i) SEQ ID NO:11;
    - ii) SEQ ID NO:12;
    - iii) SEQ ID NO:13;
    - iv) SEQ ID NO:14;
    - v) SEQ ID NO:15 after His<sup>796</sup> (SEQ ID NO:90); and
    - vi) allelic variants of any of subparts i) through v);

wherein the numbering in subpart a) is based on the amino acid sequence of SEQ ID NO:55

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9. The soluble leptin receptor of Claim 8 which is selected from the group consisting of

- C2
- a) OB-Re (SEQ ID NO:10);
  - b) an N-terminal sequence which is selected from the group consisting of:
    - i) OB-Ra (SEQ ID NO:2),
    - ii) OB-Rb (SEQ ID NO:4),
    - iii) OB-Rd (SEQ ID NO:8), and

iv) corresponding to SEQ ID NO:55 from Pro<sup>664</sup>, through His<sup>796</sup>, and a C-terminal sequence which is OB-Re from His<sup>796</sup> (SEQ ID NO:91); and

v) allelic variants of any of subparts i) through iv);

c) an N-terminal sequence which is selected from the group consisting of

i) amino acid residues 1-796 (SEQ ID NO:85);

ii) amino acid residues 23-796 (SEQ ID NO:86);

iii) amino acid residues 28-796 (SEQ ID NO:87);

iv) amino acid residues 133-796 (SEQ ID NO:88);

v) amino acid residues 733-796 (SEQ ID NO:89); and

vi) allelic variants of any of subparts i) through v); and

a C-terminal sequence which is SEQ ID NO:15;

wherein the numbering in subparts b) and c) is based on the amino acid sequence of SEQ ID NO:55.

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